SEQUENCE_Stratagene-2462.txt
SEQUENCE LISTING

<110>   Stratagene California
         Braman, Jeffrey
         Carstens, Carsten-Peter
         Novoradovskaya, Natalia
         Bagga, Rajesh
         Basehore, Lee Scott

<120>   Compositions and Methods for Protein Isolation

<130>   25436/2462

<140>   US 10/712,137
<141>   2003-11-13

<160>   32

<170>   PatentIn version 3.2

<210>   1
<211>   17
<212>   PRT
<213>   Bos taurus

<400>   1

Lys Ile Gln Ala Ser Phe Arg Gly His Ile Thr Arg Lys Lys Leu Lys
1   5   10   15

Gly

<210>   2
<211>   28
<212>   PRT
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<220>
<223>   calmodulin-dependent protein kinase I (CMKI) AA 294-318

<400>   2

Ser Glu Gln Ile Lys Lys Asn Phe Ala Lys Ser Lys Trp Lys Gln Ala
1   5   10   15

Phe Asn Ala Thr Ala Val Val Arg His Met Arg Lys
            20   25

<210>   3
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<213>   Artificial sequence

<220>
<223>   calmodulin-dependent protein kinase II (CMKII) AA 290-309

<400>   3

Leu Lys Lys Phe Asn Ala Arg Arg Lys Leu Lys Gly Ala Ile Leu Thr
1   5   10   15

Thr Met Leu Ala
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<210> 4
<211> 16
<212> PRT
<213> Homo sapiens

<400> 4

Trp Ile Ala Arg Leu Arg His Ile Lys Arg Leu Arg Gln Arg Ile Cys
1   5   10   15

<210> 5
<211> 260
<212> DNA
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<220>
<223> DNA sequence coding CBP/SBP tandem affinity tags fused to the
      N-terminus of the bait protein

<400> 5
gcggccgcca ccatgaagcg acgatggaaa aagaatttca tagccgtctc agcagccaac   60

cgctttaaga aaatctcatc ctccggggca cttggaagcg gtagcggtac catggacgag   120

aagaccaccg gctggcgggg cggccacgtg gtggagggcc tggccggcga gctggagcag   180

ctgcgggcca ggctggagca ccaccctcag ggccagcggg agccctccgg cggctgcaag   240

ctgggcgccc gggcggatcc   260

<210> 6
<211> 78
<212> PRT
<213> Artificial sequence

<220>
<223> CBP/SBP tandem affinity tags fused to the N-terminus of the bait
      protein

<400> 6

Met Lys Arg Arg Trp Lys Lys Asn Phe Ile Ala Val Ser Ala Ala Asn
1   5   10   15

Arg Phe Lys Lys Ile Ser Ser Ser Gly Ala Leu Gly Ser Gly Ser Gly
            20   25   30

Ser Met Asp Glu Lys Thr Thr Gly Trp Arg Gly Gly His Val Val Glu
        35   40   45

```
Gly Leu Ala Gly Glu Leu Glu Gln Leu Arg Ala Arg Leu Glu His His
    50                  55                  60


Pro Gln Gly Gln Arg Glu Pro Ser Gly Gly Cys Lys Leu Gly
65                  70                  75
```

<210> 7
<211> 254
<212> DNA
<213> Artificial sequence

<220>
<223> DNA sequence coding CBP/SBP tandem affinity tags fused to the
      C-terminus of the bait protein

<400> 7

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gtggtggagg gcctggccgg cgagctggag cagctgcggg ccaggctgga gcaccaccct     120

cagggccagc gggagccctc cggcggctgc aagctgggct ccggaaagcg acgatggaaa     180

aagaatttca tagccgtctc agcagccaac cgctttaaga aaatctcatc ctccggggca     240

ctttagggcc cgac                                                       254
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<210> 8
<211> 79
<212> PRT
<213> Artificial sequence

<220>
<223> CBP/SBP tandem affinity tag fused to the C-terminus of the bait
      protein

<400> 8

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Gly Ser Gly Ser Gly Ser Met Asp Glu Lys Thr Thr Gly Trp Arg Gly
1               5                  10                  15


Gly His Val Val Glu Gly Leu Ala Gly Glu Leu Glu Gln Leu Arg Ala
            20                  25                  30


Arg Leu Glu His His Pro Gln Gly Gln Arg Glu Pro Ser Gly Gly Cys
        35                  40                  45


Lys Leu Gly Ser Gly Lys Arg Arg Trp Lys Lys Asn Phe Ile Ala Val
    50                  55                  60


Ser Ala Ala Asn Arg Phe Lys Lys Ile Ser Ser Ser Gly Ala Leu
65                  70                  75
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<210> 9
<211> 101
<212> PRT

<213> Artificial sequence

<220>
<223> SBP sequence (SB1)

<400> 9

Met Asp Glu Lys Thr His Cys Thr Ile Ser Met Asn Gly Ala Val Pro
1   5   10   15

Leu Val Pro His His His Pro Gln Gly Asp Pro Leu Arg Leu Leu His
                20   25   30

Arg Pro Gln Pro Ala Leu Leu Val Arg His Pro Gln Gly Asp Leu Val
            35   40   45

Ala Leu Val Glu His His Glu Gly Val Asp Arg Gly Leu Val·Ala Leu
        50   55   60

Pro Glu Leu His Ala Glu Glu Leu Gly Glu Pro Val Gly Asp Leu Val
65   70   75   80

Gln Gly Pro Val Glu Gln Val Gln Gly Val Val Asp Ala Leu Val Trp
            85   90   95

Arg Leu Pro Pro Ser
            100

<210> 10
<211> 101
<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB2)

<400> 10

Met Asp Glu Lys Thr His Cys Phe His Pro Gly Asp His Leu Val Arg
1   5   10   15

Leu Val Glu Glu Leu Gln Ala Leu Ala Glu Gly Leu Gln Arg Gln Gly
                20   25   30

Gly Arg Gln Pro His Arg Leu Pro Arg Arg Arg Pro His His Leu Gln
            35   40   45

Leu Leu Leu Asp Glu Ala His Pro Gln Ala Gly Pro Leu Arg Glu Arg
        50   55   60

Ala His Gln Val Asp Gly Arg Leu Leu Leu Gln His His Pro Gln Gly
65   70   75   80

Asp Arg Leu Leu Gln Gln Pro Gln Asp His Pro Leu Glu Leu Val Trp
                85   90   95

Arg Leu Pro Pro Ser
            100

<210> 11
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<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB3)

<400> 11

Met Thr Arg Arg Pro Thr Ala Ser Ser Ser Ser Cys Val Arg His Leu
1   5   10   15

Leu Leu Arg Gln Gly Glu His Gly His Gln Ala Leu Glu Asp Arg Asp
                20   25   30

Lys Ala Arg His Val Arg Leu Val Glu Gly Asp Val Glu Val Leu Gly
                35   40   45

Gly Leu Asp Arg Leu Ala Arg Ala Arg His Glu Ala Leu His Pro Gln
    50   55   60

Ala Gly Leu Val His Leu Pro Leu His Gly Gly Asp Leu Gly Gly His
65   70   75   80

Leu Arg Leu Val Leu Glu Ala His Pro Gln Gly Asp Arg Leu Gly Leu
                85   90   95

Ala Val His His His
            100

<210> 12
<211> 102
<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB4)

<400> 12

Met Asp Glu Lys Thr His Trp Gly Ile Ser Thr Trp Arg Gly Glu Pro
1   5   10   15

Leu Leu His His Pro Gln Ala Gly Arg Leu Pro Leu Asp Arg Arg Arg

Ala Arg His Arg Arg Ile Leu Gly Ala Glu Pro Gly Gly Val Asp His
        35   40   45

Gly Leu Arg Leu Glu Leu Leu Asp Asp His Arg Pro Leu Val Pro Asp
        50   55   60

His His Pro Gln Arg Gly Pro Leu Gln Arg Gly Asp Leu Pro Gln Val
65   70   75   80

Val Pro Leu Val Arg Leu Arg His Ala His Val Leu Gly Leu Gly Leu
                85   90   95

Ala Ala Ala Thr Ile Thr
            100

<210>   13
<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB5)

<400>   13

Met Asp Glu Lys Thr His Trp Val Asn Val Tyr His Pro Gln Gly Asp
1   5   10   15

Leu Leu Val Arg Gly His Gly His Asp Val Glu Ala Leu His Asp Gln
                20   25   30

Gly Leu His Gln Leu Asp Leu Leu Val Gly Pro Pro Pro Glu Val Val
        35   40   45

Arg Ala Leu Arg Gly Glu Val Leu Gly Gly Leu Arg Arg Leu Val Pro
    50   55   60

Leu Asp His Pro Gln Gly Glu Ala Leu Asp Gln Ala Arg Gln Arg Pro
65   70   75   80

Gln His Leu Leu Glu Leu His His Arg Ala Leu Pro Pro Ala Leu Val
                85   90   95

Trp Arg Leu Pro Pro Ser
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<210>   14
<211>   102

<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB6)

<400> 14

Met Asp Glu Lys Thr His Trp Leu Asn Asn Phe Glu Glu Leu Leu Ala
1   5   10   15

Arg Leu Asp Gly Leu Arg Glu Gly Glu Asp His Pro Leu Val Leu Arg
                20   25   30

His His Pro Gln Gly Asp Gly Leu Leu Asp Gln Pro Leu Gly Arg His
            35   40   45

Arg Ala Leu Asp Gly Glu Val Arg Glu Gly Asp Arg Pro Leu Asp Gln
    50   55   60

Gly Gly Glu Glu Asp Leu Gly Ala Leu Val Asp Asp Asp Gly Glu Val
65   70   75   80

Leu Asp Gly Leu Val His Val Gly Val His Val His Asp Pro Leu Val
                85   90   95

Cys Gly Cys His His His
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<210> 15
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<212> PRT
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<220>
<223> SBP sequence (SB7)

<400> 15

Met Asp Glu Lys Thr His Trp Phe Gly Thr Leu Asn Ser Phe Pro Thr
1   5   10   15

His Trp Met Ser Ala Val Gly Asn Gly Lys Ile Asp Cys Ser Phe Asn
                20   25   30

Met Asn Leu Ser Leu Asn His Trp Leu Ser Ser Gly His Pro Asp Gly
            35   40   45

Ala Leu Asp Asp Gln Leu His Pro Gln Gly Asp Ala Leu Val Gly Arg
    50   55   60

Asp Asp Gly Val Val Gln Ala Leu Arg Leu Glu Gly Gln His Gln His

65   70   75   80

Arg Arg Leu Ala Gln Arg Arg Ala Asp Arg His Arg Gln Leu Val Trp
                85   90   95

Arg Leu Pro Pro Ser
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<210>   16
<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB8)

<400>   16

Met Asp Glu Lys Thr His Cys Thr Ile Glu Leu Asn Phe Ser Phe Thr
1   5   10   15

His Trp Lys Leu His His His Pro Gln Gly Asp Ala Leu Leu Asp Asp
                20   25   30

Gly Val Arg Pro His His Pro Leu Ala Asp Glu Gly Gly Gly Leu Asp
                35   40   45

Gln Gly Leu Gly His Arg Arg Gly Val Val Ala Glu Arg Leu Ala Arg
        50   55   60

Arg Asp Pro Glu Val Leu Glu Gly Leu Val Glu Arg His Arg Gly Leu
65   70   75   80

Val Pro Arg Leu Arg His Gly Gly Glu Arg His Ala Glu Pro Leu Val
                85   90   95

Trp Arg Leu Pro Pro Ser
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<210>   17
<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB9)

<400>   17

Met Asp Glu Lys Thr His Cys Asn Thr Gly Leu Tyr Asp Gly Ala Ala
1   5   10   15

```
Asp Cys Phe Asn Glu Leu Asn Lys Asp Val Ala Pro Leu Val Glu Gly
            20              25                  30


Arg His Asp Leu Val Glu Gly Leu Leu Leu Glu Arg His Pro Gln Gly
            35              40                  45


Asp Pro Leu Val Ala His Arg Gln Leu Val His His Pro Leu Leu Gly
        50              55                  60


Arg Gly Glu Arg His Arg Arg Ala Leu Val Pro Gln Gln Glu His Gln
65                  70                  75                  80


Pro His Arg Leu Gln Pro Val Val Asp Leu Gly Arg Arg Arg Leu Val
                85                  90                  95


Trp Arg Leu Pro Pro Ser
            100
```

<210> 18
<211> 103
<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB10)

<400> 18

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Met Asp Glu Lys Thr His Trp His Glu Arg Ala Gln Glu Leu Val Gly
1               5                   10                  15


Gly Leu Leu Leu His Asp His Pro Gln Arg Leu Leu Leu Glu Pro Arg
            20                  25                  30


Gly Pro Arg Pro Leu Arg Gly Leu Val His Glu Arg Gly His Gln Pro
            35              40                  45


Gln Pro Leu Ala Gly Arg Val Glu Glu Ala Asp Gly Gly Leu Leu Arg
        50              55                  60


Asp Gly Gly Gly Glu Leu Glu Pro Leu Val Arg Glu Gly Glu Asp His
65                  70                  75                  80


Leu Glu Pro Leu Asp Asp Glu Leu Asp Ala Gly Pro Arg Gly Leu Val
                85                  90                  95


Trp Arg Leu Pro His His His
            100
```

<210> 19

<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB11)

<400>   19

Met Asp Glu Lys Thr His Trp His Glu Arg Val His His Leu Ala Asp
1   5   10   15

Gly Leu Glu Gln His Pro Gln Gly Gln Arg Arg Pro Leu Val Glu Arg
                20   25   30

His Arg Gln Val Pro Arg Gly Leu Val Arg Glu Leu Gln His Glu Gly
                35   40   45

Leu Pro Leu Glu His Pro Ala Gly Val His Val Ile Arg Leu His Gln
        50   55   60

Gly Asp Asp Arg Asp Val Asp Gly Leu Val Asp Gly His Gly Arg Asp
65   70   75   80

Val Arg Gly Leu Glu Arg Glu Val Gly Asp Gly Pro His Arg Leu Val
                85   90   95

Trp Arg Leu Pro Pro Ser
                100

<210>   20
<211>   101
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB12)

<400>   20

Met Asp Lys Asp Pro Leu Leu Glu Glu Leu Glu Glu Leu Arg Glu Arg
1   5   10   15

Leu Val His His Pro Gln Gly Gly Leu Leu Pro Leu Arg Gly Gln Val
                20   25   30

Gly His Asp Ala Glu Arg Leu Gly Ala Glu Val Asp Asp Leu Arg Gly
                35   40   45

Gly Leu Leu Asp Glu Pro Gln Arg Ala Val Ala Gly Leu His His Val
        50   55   60

```
Pro His Arg Val Gly Gln Arg Leu Val His Glu Val Arg Glu Leu Asp
65              70          75              80


Glu Gly Leu Leu Asp Gln Arg Asp Asp Leu Arg Gln Arg Leu Val Trp
            85          90              95


Arg Leu Pro Pro Ser
            100
```

```
<210>   21
<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB13)

<400>   21
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```
Met Glu Arg Glu Asp Pro Leu Asp Glu Gln Leu Arg Glu Leu Arg Glu
1               5               10              15


Ala Leu Val Asp His Pro Gln Gly Gly Ala Gln Ala Leu His Arg His
            20              25              30


Asp Gly Gly Glu His Val Pro Leu Arg Arg Val Gln His Arg Leu Gln
        35              40              45


Pro Gly Leu Gln His His Leu Glu Pro Gln Pro Leu Gly Leu Leu Gly
    50              55              60


Glu Leu Gln Ala Arg Leu Gln Pro Leu Ala Gly Glu His Glu Gly Asp
65              70              75              80


Gly Ala Gly Leu Gln Arg Val Pro Gly His Gln Gly Arg Arg Leu Val
            85              90              95


Trp Arg Leu Pro Pro Ser
            100
```

```
<210>   22
<211>   101
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB14)

<400>   22
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```
Met Asp Glu Lys Thr His Arg Thr Leu Ser Val Ser Leu Ser Phe Asn
1               5               10              15
```

```
Asp Trp Leu Gly Gln Thr Lys Ala Cys Trp Arg Leu Val Glu Gly Leu
            20              25              30

His Gly His Pro Gln Gly Leu Val Arg Glu His Glu Val Asp Val Leu
            35              40              45

Pro Leu Ala Glu Glu Val Gln Gln Val Val Gly Gly Leu Ala Asp Gly
        50              55              60

Val Glu Gln Pro Gly Gly Gly Leu Leu His Arg Ala Gln Arg Val Asp
65              70              75              80

His Pro Leu Pro Asp His Ala Gly Gln Val Leu Gly Arg Leu Val Trp
                85              90              95

Arg Leu Pro Pro Ser
            100
```

<210> 23
<211> 101
<212> PRT
<213> Artificial sequence

<220>
<223> SBP sequence (SB15)

<400> 23

```
Met Asp Glu Lys Thr His Trp Leu Glu Asp Leu Lys Gly Val Leu Lys
1               5               10              15

Asp Cys Leu Lys Asp Leu Met Asp Phe Thr Lys Asp Cys Arg Ser Pro
            20              25              30

Arg Val Gln Pro Gln Pro Leu Leu His His Asp Arg Gly Glu Pro Val
            35              40              45

Pro Leu Leu Arg Glu Ala Gly Arg Asp Leu Gly Gly Leu Gly Pro Arg
        50              55              60

Ala Pro Arg Gln Ala Arg Pro Leu His His Gly Arg His Asp Leu His
65              70              75              80

Glu Pro Leu Val Leu Gln Asp His Pro Gln Gly Gly Pro Leu Val Cys
                85              90              95

Gly Cys His His His
            100
```

```
<210>  24
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<213>  Artificial sequence

<220>
<223>  SBP sequence (SB16)

<400>  24
```

Met Asp Glu Lys Thr His Trp Val Leu Gln Leu His Pro Gln Gly Asp
1   5   10   15

Arg Leu Gly Pro Arg His Gly Gly Asp Asp Val Arg Leu Val Gly Gln
            20   25   30

Gly Glu Gly Val Leu Glu Gly Leu Asp Gly Arg Pro Arg Arg Arg Arg
            35   40   45

His Arg Leu Pro Arg Glu Asp Glu His Arg Val Arg Ala Leu Val Asp
        50   55   60

Gln Val Arg Asp Leu Ala Glu Arg Leu Val Glu Glu Val Asp Gly Gly
65   70   75   80

Val Glu Ala Leu Arg His Leu Gly Leu Pro Gln Asp Glu Pro Arg Ser
                85   90   95

Gly Gly Cys His His His
                100

```
<210>  25
<211>  102
<212>  PRT
<213>  Artificial sequence

<220>
<223>  SBP sequence (SB17)

<400>  25
```

Met Asp Glu Lys Thr His Trp Val Gly Asp Leu Gln Glu Pro Leu Gly
1   5   10   15

Pro Leu His Gly Gly Val Gly Glu Val Pro Gly Gly Leu Val Leu Arg
            20   25   30

His His Pro Gln Arg Asp Arg Leu Val Asp Gly Val Gly Pro His Gly
            35   40   45

Arg Ala Leu Ala Arg Arg Pro His Arg Val Val Glu Gly Leu His His
        50   55   60

Leu Leu Gln Arg Gly Gly Glu Arg Leu Pro Pro Asp Gly Pro Arg Gln
65   70   75   80

Leu Gly Leu Leu Gly Gly Glu Leu Asp Arg Ala Asp Pro Ala Leu Val
                85   90   95

Trp Arg Leu Pro Pro Ser
                100

<210>   26
<211>   101
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB18)

<400>   26

Met Asp Glu Lys Thr His Cys Ala Val Asn Val Asn Val Gly Leu Thr
1   5   10   15

His Trp Cys His Arg Val Ala His Leu Gln Pro Leu Asp Pro His Pro
                20   25   30

Gln Gly Asp His Leu Arg Leu Glu Pro Leu Gly His Ala Leu Val Asp
        35   40   45

Pro Leu Val Gln Gly Val Glu Glu Val Val Arg Pro Leu Gln Leu Asp
    50   55   60

Val Gly Val Gln Arg Val Ala Leu Val Glu Gln Val Ala Glu Val Gly
65   70   75   80

Glu Gly Leu Asp His Glu Ala Gly Gln Ala His Gly Ala Leu Val Trp
                85   90   95

Arg Leu Pro Pro Ser
                100

<210>   27
<211>   101
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB19)

<400>   27

Met Asp Glu Lys Thr Thr Gly Trp Arg Gly Gly His Val Val Glu Gly
1   5   10   15

```
Leu Ala Gly Glu Leu Glu Gln Leu Arg Ala Arg Leu Glu His His Pro
            20              25              30

Gln Gly Gln Arg Glu Pro Leu Val Gln Glu Val Glu Asp Val Asp Glu
            35              40              45

Gly Leu Val Gln Asp Leu His Gly Val Val Ala Gly Leu Leu Asp Pro
        50              55              60

Val Glu Lys Leu Leu Thr Asp Trp Phe Lys Lys Phe Lys Asn Val Ser
65                  70              75              80

Lys Asp Cys Lys Met Thr Phe Tyr Leu Glu Met Tyr Asp Trp Ser Gly
                85              90              95

Gly Cys His His His
            100
```

<210>   28
<211>   102
<212>   PRT
<213>   Artificial sequence

<220>
<223>   SBP sequence (SB20)

<400>   28

```
Met Asn Glu Lys Thr His Cys Lys Leu Asn Phe Lys Val Asn Ile Ala
1               5               10              15

Asp Trp Leu Ala Glu Phe His Gly Gly Gly Gln Gly Leu Leu Gly Arg
            20              25              30

Arg Asp Gly Val Val Gln Arg Leu Val Asp Gly Val Gln Glu Arg Val
            35              40              45

Glu Arg Leu Asp Arg Asp Pro Gly Leu Gly Asp Leu Arg Leu Glu Leu
        50              55              60

His His Arg Asp His Arg Leu Arg Leu Gly Gly Glu His Leu Leu Arg
65                  70              75              80

Asp His Pro Leu Glu Pro Asp Asp His Leu Val Val Gly Gly Leu Val
                85              90              95

Trp Arg Leu Pro Pro Ser
            100
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<210> 29
<211> 4531
<212> DNA
<213> Artificial sequence

<220>
<223> Expression vector comprising nucleic acids encoding CBP and SBP
affinity tags

<400> 29

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cccattgacg tcaataatga cgtatgttcc catagtaacg ccaataggga ctttccattg      180

acgtcaatgg gtggagtatt tacggtaaac tgcccacttg gcagtacatc aagtgtatca      240

tatgccaagt acgcccccta ttgacgtcaa tgacggtaaa tggcccgcct ggcattatgc      300

ccagtacatg accttatggg actttcctac ttggcagtac atctacgtat tagtcatcgc      360

tattaccatg gtgatgcggt tttggcagta catcaatggg cgtggatagc ggtttgactc      420

acggggattt ccaagtctcc accccattga cgtcaatggg agtttgtttt ggcaccaaaa      480

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gcgattacgc caagctcgaa attaaccctc actaaaggga acaaaagctg gagctccacc      660

gcggtggcgg ccgccaccat gaagcgacga tggaaaaaga atttcatagc cgtctcagca      720

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tacccaattc gccctatagt gagtcgtatt acaattcact cgatcgccct tcccaacagt     1080

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tcctcacagt ctgttcatga tcataatcag ccataccaca tttgtagagg ttttacttgc     1260

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cacaaataaa gcattttttt cactgcattc tagttgtggt ttgtccaaac tcatcaatgt     1440

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aaatcagctc attttttaac caataggccg aaatcggcaa aatcccttat aaatcaaaag     1560
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ctgaggcgga aagaaccagc tgtggaatgt gtgtcagtta gggtgtggaa agtccccagg   2100

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cttttttacgg ttcctggcct tttgctggcc ttttgctcac atgttctttc ctgcgttatc    4500

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<212> DNA
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<400> 30
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cccattgacg tcaataatga cgtatgttcc catagtaacg ccaatagggga ctttccattg    180

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<213>  Artificial sequence

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cccattgacg tcaataatga cgtatgttcc catagtaacg ccaatagga ctttccattg     180

acgtcaatgg gtggagtatt tacggtaaac tgcccacttg gcagtacatc aagtgtatca    240

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